RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/552,061_

Source: _PCT/10_

Date Processed by STIC: _10/17/05_

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PCT

RAW SEQUENCE LISTING   DATE: 10/17/2005
PATENT APPLICATION: US/10/552,061   TIME: 10:57:43

Input Set : A:\5.1277.1.txt
Output Set: N:\CRF4\10172005\J552061.raw

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  3 <110> APPLICANT: KYOWA HAKKO KOGYO CO., LTD.
  5 <120> TITLE OF INVENTION: Nerve Regenerating Drug
  7 <130> FILE REFERENCE: 5.1277
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/552,061
C--> 10 <141> CURRENT FILING DATE: 2005-10-04
 12 <150> PRIOR APPLICATION NUMBER: PCT/JP04/005503
 13 <151> PRIOR FILING DATE: 2004-04-16
 15 <150> PRIOR APPLICATION NUMBER: JP2003-114579
 16 <151> PRIOR FILING DATE: 2003-04-18
 18 <160> NUMBER OF SEQ ID NOS: 17
 20 <170> SOFTWARE: PatentIn Ver. 2.1
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 23 <211> LENGTH: 420
 24 <212> TYPE: PRT
 25 <213> ORGANISM: Homo sapiens
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 34 Asp Gly Ser Lys Val Thr Thr Val Val Ala Thr Pro Gly Gln Gly Pro
 35             35                  40                  45
 37 Asp Arg Pro Gln Glu Val Ser Tyr Thr Asp Thr Lys Val Ile Gly Asn
 38         50                  55                  60
 40 Gly Ser Phe Gly Val Val Tyr Gln Ala Lys Leu Cys Asp Ser Gly Glu
 41 65                  70                  75                  80
 43 Leu Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Phe Lys Asn Arg
 44                 85                  90                  95
 46 Glu Leu Gln Ile Met Arg Lys Leu Asp His Cys Asn Ile Val Arg Leu
 47                 100                 105                 110
 49 Arg Tyr Phe Phe Tyr Ser Ser Gly Glu Lys Lys Asp Glu Val Tyr Leu
 50             115                 120                 125
 52 Asn Leu Val Leu Asp Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg
 53         130                 135                 140
 55 His Tyr Ser Arg Ala Lys Gln Thr Leu Pro Val Ile Tyr Val Lys Leu
 56 145                 150                 155                 160
 58 Tyr Met Tyr Gln Leu Phe Arg Ser Leu Ala Tyr Ile His Ser Phe Gly
 59                 165                 170                 175
 61 Ile Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Leu Asp Pro Asp
 62             180                 185                 190
 64 Thr Ala Val Leu Lys Leu Cys Asp Phe Gly Ser Ala Lys Gln Leu Val
 65             195                 200                 205
 67 Arg Gly Glu Pro Asn Val Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala
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71 225                 230                 235                 240
73 Trp Ser Ala Gly Cys Val Leu Ala Glu Leu Leu Leu Gly Gln Pro Ile
74              245                 250                 255
76 Phe Pro Gly Asp Ser Gly Val Asp Gln Leu Val Glu Ile Ile Lys Val
77              260                 265                 270
79 Leu Gly Thr Pro Thr Arg Glu Gln Ile Arg Glu Met Asn Pro Asn Tyr
80          275                 280                 285
82 Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp Thr Lys Val
83      290                 295                 300
85 Phe Arg Pro Arg Thr Pro Pro Glu Ala Ile Ala Leu Cys Ser Arg Leu
86 305                 310                 315                 320
88 Leu Glu Tyr Thr Pro Thr Ala Arg Leu Thr Pro Leu Glu Ala Cys Ala
89              325                 330                 335
91 His Ser Phe Phe Asp Glu Leu Arg Asp Pro Asn Val Lys His Pro Asn
92          340                 345                 350
94 Gly Arg Asp Thr Pro Ala Leu Phe Asn Phe Thr Thr Gln Glu Leu Ser
95          355                 360                 365
97 Ser Asn Pro Pro Leu Ala Thr Ile Leu Ile Pro Pro His Ala Arg Ile
98      370                 375                 380
100 Gln Ala Ala Ala Ser Thr Pro Thr Asn Ala Thr Ala Ala Ser Asp Ala
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123 gac ggc agc aag gtg aca aca gtg gtg gca act cct ggg cag ggt cca   144
124 Asp Gly Ser Lys Val Thr Thr Val Val Ala Thr Pro Gly Gln Gly Pro
125          35                  40                  45
127 gac agg cca caa gaa gtc agc tat aca gac act aaa gtg att gga aat   192
128 Asp Arg Pro Gln Glu Val Ser Tyr Thr Asp Thr Lys Val Ile Gly Asn
129      50                  55                  60
131 gga tca ttt ggt gtg gta tat caa gcc aaa ctt tgt gat tca gga gaa   240
132 Gly Ser Phe Gly Val Val Tyr Gln Ala Lys Leu Cys Asp Ser Gly Glu
133 65                  70                  75                  80
135 ctg gtc gcc atc aag aaa gta ttg cag gac aag aga ttt aag aat cga   288
136 Leu Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Phe Lys Asn Arg
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144 Arg Tyr Phe Phe Tyr Ser Ser Gly Glu Lys Lys Asp Glu Val Tyr Leu
145          115                     120                  125
147 aat ctg gtg ctg gac tat gtt ccg gaa aca gta tac aga gtt gcc aga    432
148 Asn Leu Val Leu Asp Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg
149      130                     135                  140
151 cac tat agt cga gcc aaa cag acg ctc cct gtg att tat gtc aag ttg    480
152 His Tyr Ser Arg Ala Lys Gln Thr Leu Pro Val Ile Tyr Val Lys Leu
153 145                     150                     155              160
155 tat atg tat cag ctg ttc cga agt tta gcc tat atc cat tcc ttt gga    528
156 Tyr Met Tyr Gln Leu Phe Arg Ser Leu Ala Tyr Ile His Ser Phe Gly
157                  165                     170                  175
159 atc tgc cat cgg gat att aaa ccg cag aac ctc ttg ttg gat cct gat    576
160 Ile Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Leu Asp Pro Asp
161              180                     185                  190
163 act gct gta tta aaa ctc tgt gac ttt gga agt gca aag cag ctg gtc    624
164 Thr Ala Val Leu Lys Leu Cys Asp Phe Gly Ser Ala Lys Gln Leu Val
165          195                     200                  205
167 cga gga gaa ccc aat gtt tcg tat atc tgt tct cgg tac tat agg gca    672
168 Arg Gly Glu Pro Asn Val Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala
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175 tgg tct gct ggc tgt gtg ttg gct gag ctg tta cta gga caa cca ata    768
176 Trp Ser Ala Gly Cys Val Leu Ala Glu Leu Leu Leu Gly Gln Pro Ile
177                  245                     250                  255
179 ttt cca ggg gat agt ggt gtg gat cag ttg gta gaa ata atc aag gtc    816
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184 Leu Gly Thr Pro Thr Arg Glu Gln Ile Arg Glu Met Asn Pro Asn Tyr
185          275                     ·280                  285
187 aca gaa ttt aaa ttc cct caa att aag gca cat cct tgg act aag gtc    912
188 Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp Thr Lys Val
189      290                     295                  300
191 ttc cga ccc cga act cca ccg gag gca att gca ctg tgt agc cgt ctg    960
192 Phe Arg Pro Arg Thr Pro Pro Glu Ala Ile Ala Leu Cys Ser Arg Leu
193 305                     310                     315              320
195 ctg gag tat aca cca act gcc cga cta aca cca ctg gaa gct tgt gca    1008
196 Leu Glu Tyr Thr Pro Thr Ala Arg Leu Thr Pro Leu Glu Ala Cys Ala
197                  325                     330                  335
199 cat tca ttt ttt gat gaa tta cgg gac cca aat gtc aaa cat cca aat    1056
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207 agt aat cca cct ctg gct acc atc ctt att cct cct cat gct cgg att    1152
208 Ser Asn Pro Pro Leu Ala Thr Ile Leu Ile Pro Pro His Ala Arg Ile
209     370                 375                 380
211 caa gca gct gct tca acc ccc aca aat gcc aca gca gcg tca gat gct    1200
212 Gln Ala Ala Ala Ser Thr Pro Thr Asn Ala Thr Ala Ala Ser Asp Ala
213 385                 390                 395                 400
215 aat act gga gac cgt gga cag acc aat aat gct gct tct gca tca gct    1248
216 Asn Thr Gly Asp Arg Gly Gln Thr Asn Asn Ala Ala Ser Ala Ser Ala
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220 Ser Asn Ser Thr
221             420
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224 <211> LENGTH: 737
225 <212> TYPE: PRT
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233             20                  25                  30
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236         35                  40                  45
238 Thr Lys Ala Lys Val Thr Gly Asp Glu Trp Gly Asp Asn Tyr Phe Leu
239     50                  55                  60
241 Val Gly Pro Tyr Thr Glu Gln Gly Val Arg Thr Gln Val Glu Leu Leu
242 65                  70                  75                  80
244 Glu Ala Pro Thr Pro Ala Leu Lys Arg Thr Leu Asp Ser Met Asn Ser
245             85                  90                  95
247 Lys Gly Cys Lys Val Tyr Phe Gly Arg Trp Leu Ile Glu Gly Gly Pro
248         100                 105                 110
250 Leu Val Val Leu Leu Asp Val Gly Ala Ser Ala Trp Ala Leu Glu Arg
251         115                 120                 125
253 Trp Lys Gly Glu Leu Trp Asp Ile Cys Asn Ile Gly Val Pro Trp Tyr
254 130                 135                 140
256 Asp Arg Glu Ala Asn Asp Ala Val Leu Phe Gly Phe Leu Thr Thr Trp
257 145                 150                 155                 160
259 Phe Leu Gly Glu Phe Leu Ala Gln Ser Glu Glu Lys Pro His Val Val
260                 165                 170                 175
262 Ala His Phe His Glu Trp Leu Ala Gly Val Gly Leu Cys Leu Cys Arg
263             180                 185                 190
265 Ala Arg Arg Leu Pro Val Ala Thr Ile Phe Thr Thr His Ala Thr Leu
266             195                 200                 205
268 Leu Gly Arg Tyr Leu Cys Ala Gly Ala Val Asp Phe Tyr Asn Asn Leu
269     210                 215                 220
271 Glu Asn Phe Asn Val Asp Lys Glu Ala Gly Glu Arg Gln Ile Tyr His
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272 225                    230                235                240
274 Arg Tyr Cys Met Glu Arg Ala Ala Ala His Cys Ala His Val Phe Thr
275              245                250              255
277 Thr Val Ser Gln Ile Thr Ala Ile Glu Ala Gln His Leu Leu Lys Arg
278              260                265              270
280 Lys Pro Asp Ile Val Thr Pro Asn Gly Leu Asn Val Lys Lys Phe Ser
281              275                280              285
283 Ala Met His Glu Phe Gln Asn Leu His Ala Gln Ser Lys Ala Arg Ile
284              290                295              300
286 Gln Glu Phe Val Arg Gly His Phe Tyr Gly His Leu Asp Phe Asn Leu
287 305              310                315              320
289 Asp Lys Thr Leu Tyr Phe Phe Ile Ala Gly Arg Tyr Glu Phe Ser Asn
290              325                330              335
292 Lys Gly Ala Asp Val Phe Leu Glu Ala Leu Ala Arg Leu Asn Tyr Leu
293              340                345              350
295 Leu Arg Val Asn Gly Ser Glu Gln Thr Val Val Ala Phe Phe Ile Met
296              355                360              365
298 Pro Ala Arg Thr Asn Asn Phe Asn Val Glu Thr Leu Lys Gly Gln Ala
299              370                375              380
301 Val Arg Lys Gln Leu Trp Asp Thr Ala Asn Thr Val Lys Glu Lys Phe
302 385              390                395              400
304 Gly Arg Lys Leu Tyr Glu Ser Leu Leu Val Gly Ser Leu Pro Asp Met
305              405                410              415
307 Asn Lys Met Leu Asp Lys Glu Asp Phe Thr Met Met Lys Arg Ala Ile
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310 Phe Ala Thr Gln Arg Gln Ser Phe Pro Pro Val Cys Thr His Asn Met
311              435                440              445
313 Leu Asp Asp Ser Ser Asp Pro Ile Leu Thr Thr Ile Arg Arg Ile Gly
314              450                455              460
316 Leu Phe Asn Ser Ser Ala Asp Arg Val Lys Val Ile Phe His Pro Glu
317 465              470                475              480
319 Phe Leu Ser Ser Thr Ser Pro Leu Leu Pro Val Asp Tyr Glu Glu Phe
320              485                490              495
322 Val Arg Gly Cys His Leu Gly Val Phe Pro Ser Tyr Tyr Glu Pro Trp
323              500                505              510
325 Gly Tyr Thr Pro Ala Glu Cys Thr Val Met Gly Ile Pro Ser Ile Ser
326              515                520              525
328 Thr Asn Leu Ser Gly Phe Gly Cys Phe Met Glu Glu His Ile Ala Asp
329              530                535              540
331 Pro Ser Ala Tyr Gly Ile Tyr Ile Leu Asp Arg Arg Phe Arg Ser Leu
332 545              550                555              560
334 Asp Asp Ser Cys Ser Gln Leu Thr Ser Phe Leu Tyr Ser Phe Cys Gln
335              565                570              575
337 Gln Ser Arg Arg Gln Arg Ile Ile Gln Arg Asn Arg Thr Glu Arg Leu
338              580                585              590
340 Ser Asp Leu Leu Asp Trp Lys Tyr Leu Gly Arg Tyr Tyr Met Ser Ala
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PATENT APPLICATION: US/10/552,061　　　TIME: 10:57:44

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